#2 OIPE

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> This Raw Listing contains the General
> Information Section and up to the first 5 pages.

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1                              SEQUENCE LISTING
2
3   (1)    General Information:
4
5       (i) APPLICANT: HOSTETTER, MARGARET K.
6                     GALE, CHERYL A.
7                     BENDEL, CATHERINE M.
8                     TAO, NIAN-JUN
9                     KENDRICK, KATHLEEN
10
11     (ii) TITLE OF INVENTION: CANDIDA ALBICANS GENE, INTEGRIN-LIKE
12              PROTEIN, ANTIBODIES, AND METHODS OF USE
13
14    (iii) NUMBER OF SEQUENCES: 12
15
16     (iv) CORRESPONDENCE ADDRESS:
17          (A) ADDRESSEE: MUETING, RAASCH, GEBHARDT & SCHWAPPACH, P.A.
18          (B) STREET: 119 NORTH FOURTH STREET, SUITE 203
19          (C) CITY: MINNEAPOLIS
20          (D) STATE: MINNESOTA
21          (E) COUNTRY: USA
22          (F) ZIP: 55401
23
24      (v) COMPUTER READABLE FORM:
25          (A) MEDIUM TYPE: Floppy disk
26          (B) COMPUTER: IBM PC compatible
27          (C) OPERATING SYSTEM: PC-DOS/MS-DOS
28          (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
29
30     (vi) CURRENT APPLICATION DATA:
31          (A) APPLICATION NUMBER: 09/978,343
32          (B) FILING DATE:
33          (C) CLASSIFICATION:
34
35    (vii) PRIOR APPLICATION DATA:
36          (A) APPLICATION NUMBER: US 08/642,846
37          (B) FILING DATE: 03-MAY-1996
38
39
40
41   (viii) ATTORNEY/AGENT INFORMATION:
42          (A) NAME: MUETING, ANN M.
43          (B) REGISTRATION NUMBER: 33,977
44          (C) REFERENCE/DOCKET NUMBER: 110.00280101
45
46     (ix) TELECOMMUNICATION INFORMATION:
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47   (A) TELEPHONE: 612-305-1217
48   (B) TELEFAX: 612-305-1228
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51 (2) INFORMATION FOR SEQ ID NO:1:
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53   (i) SEQUENCE CHARACTERISTICS:
54   (A) LENGTH: 5194 base pairs
55   (B) TYPE: nucleic acid
56   (C) STRANDEDNESS: single
57   (D) TOPOLOGY: linear
58
59   (ii) MOLECULE TYPE: DNA (genomic)
60
61
62
63
64   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
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RAW SEQUENCE LISTING
PATENT APPLICATION   *US/09/978,343*

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240    (2)  INFORMATION FOR SEQ ID NO:2:
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242         (i)  SEQUENCE CHARACTERISTICS:
243              (A)  LENGTH: 1664 amino acids
244              (B)  TYPE: amino acid
245              (C)  STRANDEDNESS: single
246              (D)  TOPOLOGY: linear
247
248         (ii)  MOLECULE TYPE: protein
249
250
251
252
253         (xi)  SEQUENCE DESCRIPTION: SEQ ID NO:2:
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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION   *US/09/978,343*

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Line	Error	Original Text

SEQUENCE MISSING ITEM REPORT
PATENT APPLICATION   *US/09/978,343*

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SEQUENCE CORRECTION REPORT
PATENT APPLICATION *US/09/978,343*

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